

1743



ENTERED

1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/508,775

DATE: 11/19/2002

TIME: 10:29:47

Input Set : D:\403uspc.app.txt

Output Set: N:\CRF4\11192002\I508775.raw

RECEIVED

DEC 03 2002

TC 1700

4 <110> APPLICANT: Mattiasson, Bo
 5 Csoregi, Elisabeth
 6 Bontidean, Ibolya
 7 Johansson, Gillis
 8 Berggren, Christine
 9 Brown, Nigel
 10 Lloyd, Jonathan
 11 Jakeman, Kenneth
 12 Hobman, Jonathan
 13 Wilson, Jonathan
 14 Van Der Leile, Daniel
 15 Corbisier, Philippe
 18 <120> TITLE OF INVENTION: METAL ION SPECIFIC CAPACITY AFFINITY SENSOR
 21 <130> FILE REFERENCE: 100096.403USPC
 23 <140> CURRENT APPLICATION NUMBER: US 09/508,775
 C--> 24 <141> CURRENT FILING DATE: 2000-05-17
 26 <160> NUMBER OF SEQ ID NOS: 4
 28 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 30 <210> SEQ ID NO: 1
 31 <211> LENGTH: 289
 32 <212> TYPE: PRT
 33 <213> ORGANISM: Synechococcus sp.
 35 <400> SEQUENCE: 1
 36 Met Ser Pro Ile Leu Gly Tyr Trp Lys Ile Lys Gly Leu Val Gln Pro
 37 1 5 10 15
 38 Thr Arg Leu Leu Leu Glu Tyr Leu Glu Glu Lys Tyr Glu Glu His Leu
 39 20 25 30
 40 Tyr Glu Arg Asp Glu Gly Asp Lys Trp Arg Asn Lys Lys Phe Glu Leu
 41 35 40 45
 42 Gly Leu Glu Phe Pro Asn Leu Pro Tyr Tyr Ile Asp Gly Asp Val Lys
 43 50 55 60
 44 Leu Thr Gln Ser Met Ala Ile Ile Arg Tyr Ile Ala Asp Lys His Asn
 45 65 70 75 80
 46 Met Leu Gly Gly Cys Pro Lys Glu Arg Ala Glu Ile Ser Met Leu Glu
 47 85 90 95
 48 Gly Ala Val Leu Asp Ile Arg Tyr Gly Val Ser Arg Ile Ala Tyr Ser
 49 100 105 110
 50 Lys Asp Phe Glu Thr Leu Lys Val Asp Phe Leu Ser Lys Leu Pro Glu
 51 115 120 125
 52 Met Leu Lys Met Phe Glu Asp Arg Leu Cys His Lys Thr Tyr Leu Asn
 53 130 135 140
 54 Gly Asp His Val Thr His Pro Asp Phe Met Leu Tyr Asp Ala Leu Asp
 55 145 150 155 160

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56 Val Val Leu Tyr Met Asp Pro Met Cys Leu Asp Ala Phe Pro Lys Leu
57          165          170          175
58 Val Cys Phe Lys Lys Arg Ile Glu Ala Ile Pro Gln Ile Asp Lys Tyr
59          180          185          190
60 Leu Lys Ser Ser Lys Tyr Ile Ala Trp Pro Leu Gln Gly Trp Gln Ala
61          195          200          205
62 Thr Phe Gly Gly Gly Asp His Pro Pro Lys Ser Asp Leu Ile Glu Gly
63          210          215          220
64 Arg Gly Ile Pro Met Thr Ser Thr Thr Leu Val Lys Cys Ala Cys Glu
65 225          230          235          240
66 Pro Cys Leu Cys Asn Val Asp Pro Ser Lys Ala Ile Asp Arg Asn Gly
67          245          250          255
68 Leu Tyr Tyr Cys Ser Glu Ala Cys Ala Asp Gly His Thr Gly Gly Ser
69          260          265          270
70 Lys Gly Cys Gly His Thr Gly Cys Asn Cys Ser Glu Phe Ile Val Thr
71          275          280          285
72 Asp
75 <210> SEQ ID NO: 2
76 <211> LENGTH: 144
77 <212> TYPE: PRT
78 <213> ORGANISM: Pseudomonas aeruginosa
80 <400> SEQUENCE: 2
81 Met Glu Asn Asn Leu Glu Asn Leu Thr Ile Gly Val Phe Ala Lys Ala
82 1          5          10          15
83 Ala Gly Val Asn Val Glu Thr Ile Arg Phe Tyr Gln Arg Lys Gly Leu
84          20          25          30
85 Leu Leu Glu Pro Asp Lys Pro Tyr Gly Ser Ile Arg Arg Tyr Gly Glu
86          35          40          45
87 Ala Asp Val Thr Arg Val Arg Phe Val Lys Ser Ala Gln Arg Leu Gly
88          50          55          60
89 Phe Ser Leu Asp Glu Ile Ala Glu Leu Leu Arg Leu Glu Asp Gly Thr
90 65          70          75          80
91 His Cys Glu Glu Ala Ser Ser Leu Ala Glu His Lys Leu Lys Asp Val
92          85          90          95
93 Arg Glu Lys Met Ala Asp Leu Ala Arg Met Glu Ala Val Leu Ser Glu
94          100          105          110
95 Leu Val Cys Ala Cys His Ala Arg Arg Gly Asn Val Ser Cys Pro Leu
96          115          120          125
97 Ile Ala Ser Leu Gln Gly Gly Ala Ser Leu Ala Gly Ser Ala Met Pro
98          130          135          140
100 <210> SEQ ID NO: 3
101 <211> LENGTH: 145
102 <212> TYPE: PRT
103 <213> ORGANISM: Alcaligenes eutrophus
105 <400> SEQUENCE: 3
106 Met Asn Ile Gln Ile Gly Glu Leu Ala Lys Arg Thr Ala Cys Pro Val
107 1          5          10          15
108 Val Thr Ile Arg Phe Tyr Glu Gln Glu Gly Leu Leu Pro Pro Pro Gly
109          20          25          30

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```

110 Arg Ser Arg Gly Asn Phe Arg Leu Tyr Gly Glu Glu His Val Glu Arg
111      35      40      45
112 Leu Gln Phe Ile Arg His Cys Arg Ser Leu Asp Met Pro Leu Ser Asp
113      50      55      60
114 Val Arg Thr Leu Leu Ser Tyr Arg Lys Arg Pro Asp Gln Asp Cys Gly
115 65      70      75      80
116 Glu Val Asn Met Leu Leu Asp Glu His Ile Arg Gln Val Glu Ser Arg
117      85      90      95
118 Ile Gly Ala Leu Leu Glu Leu Lys His His Leu Val Glu Leu Arg Glu
119      100      105      110
120 Ala Cys Ser Gly Ala Arg Pro Ala Gln Ser Cys Gly Ile Leu Gln Gly
121      115      120      125
122 Leu Ser Asp Cys Val Cys Asp Thr Arg Gly Thr Thr Ala His Pro Ser
123      130      135      140
124 Asp
125 145
127 <210> SEQ ID NO: 4
128 <211> LENGTH: 72
129 <212> TYPE: PRT
130 <213> ORGANISM: Pseudomonas aeruginosa
132 <400> SEQUENCE: 4
133 Ala Thr Gln Thr Val Thr Leu Ser Val Pro Gly Met Thr Cys Ser Ala
134 1      5      10      15
135 Cys Pro Ile Thr Val Lys Lys Ala Ile Ser Glu Val Glu Gly Val Ser
136      20      25      30
137 Lys Val Asp Val Thr Phe Glu Thr Arg Gln Ala Val Val Thr Phe Asp
138      35      40      45
139 Asp Ala Lys Thr Ser Val Gln Lys Leu Thr Lys Ala Thr Ala Asp Ala
140      50      55      60
141 Gly Tyr Pro Ser Ser Val Lys Gln
142 65      70

```

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/508,775

DATE: 11/19/2002

TIME: 10:29:48

Input Set : D:\403uspc.app.txt

Output Set: N:\CRF4\11192002\I508775.raw

L:24 M:271 C: Current Filing Date differs, Replaced Current Filing Date